

1/25

Fig. 1A
SEQ. ID NO:1

MTVARPSVPAALPLLGELPRLLLVLCLPAVGDCGLPPDVPNAQPALE 50
GRTSFPEDTVITYKCEESFVKIPGEKDSVICLKGSQWSDIIEFCNRSCEV 100
PTRLNSASLKQPYITQNYFPVGTVVEYECPGXRREPSLSPKLTCQLQNLK 150
WSTAVEFCKKKSCPNPGEIRNGQIDVPGGILFGATISFSCNTGYKLFGST 200
SSFCLISGSSVQWSDPLPECREIYCPAPPQIDNGIIQGERDHGYRQSVT 250
YACNKGFTMIGEHSIYCTVNNDEGEWSGPPPECRGKSLTSKVPPTVQKPT 300
TVNVPTTEVSPTSQKTTKTTTPNAQATRSTPVSRTTKHFHETTPNKGSG 350
TTSGGTTRLLSGHTCFTLTGLLGTIVTMGLLT

Fig. 1B
SEQ. ID NO:2

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181 ccccaagatgt acctaattgcc cagccagctt tggaaaggccg tacaagttt cccgaggata
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2041 agaagatatg tgaagaaaaa aatgttgc acgttgacag gtttgcttgg gtttccagg gtttccagg ttcaggtact acccgtctc
2101 gt

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Fig. 2

SEQ. ID NO:3

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GILVSDNRSLSLNEVVEFRCQPGFVMKGPRRVKCQALNKWEPELPCSR	300
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SASYCVLAGMESLWNSSVPVCEQIFCPSPPVIPNGRHTGKPLEVFPFGKA	900
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PDHFLFAKLKTQTNASDFPIGTSLKYECRPEYYGRPFSITCLDNLVWSSP	1000
KDVCKRKSCCKTPPDVNGMVHVTIDIQVGSRINYSCTTGHRLIGHSSAEC	1050
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GSPWSQCQADDRWDPLAKCTSRAHDALIVGTLSTGIFTILLIIFLSWII	2000
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Fig. 3

SEQ. ID NO:4

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Fig. 4A

SEQ. ID NO:5

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Fig. 4B

SEQ. ID NO:6

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 1021 tttggatgtt tgggtcattt ctgtgattt tattgccata gttgttggag ttgcagtaat
 1081 ttgtgttgc ccgtacagat atcttcaaag gaggaaggaa aaaggaaaag cagatggtgg
 1141 agctgaatat gccacttacc agactaaatc aaccactcca gcagagcaga gaggctgaat
 1201 agattccaca acctggttt ccagttcatc ttttactt attaaaaatct tcaatagtt
 1261 ttattctgtt gtttacttcatc catgagtgc actgtggctt agctaattt gcaatgtggc
 1321 ttgaatgtt gtagcatcct ttgatgctt ttgaaactt gtatgaattt gggtatgaac
 1381 agattgcctg ctttccctt aataacactt agatttattt gaccagtcag cacagcatgc
 1441 ctgggtgtat taaagcaggg atatgctgtt tttataaaa ttggcaaaat tagagaaata
 1501 tagttcacaa tgaaattata ttttctttgtt

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Fig. 5

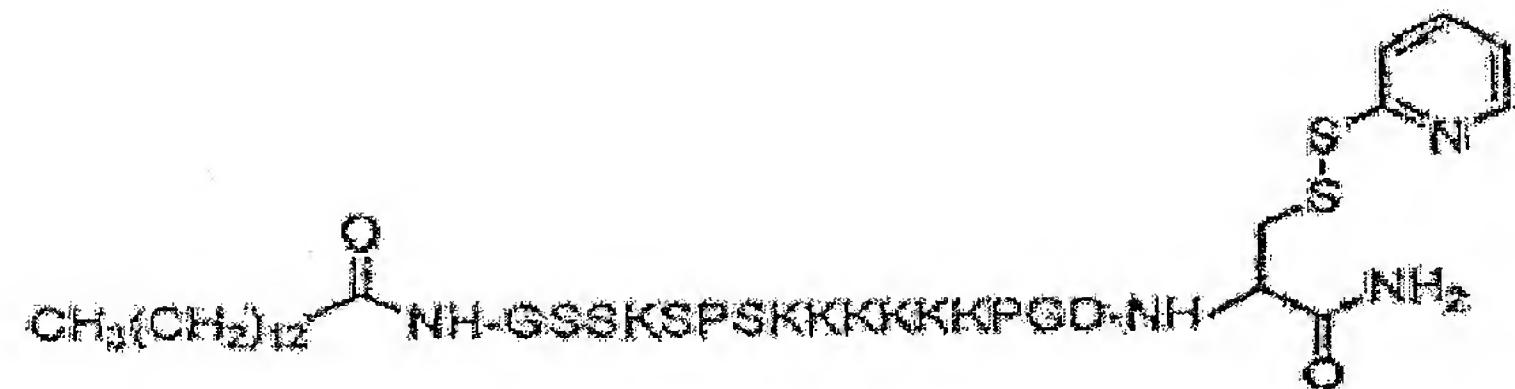


Fig. 6A

SEQ. ID NO:7

ATA TAC GAA TTC AGA TCT ATG ACC GTC GCG CGG CCG AGC GTG

Fig. 6B

SEQ. ID NO:8

ACA GTG CTC GAG CAT TCA GGT GGT GGG CCA CTC CA

Fig 7A

SEQ. ID NO:9

ATA TAC CTC GAG TCC TAA CAA ATG CAC GCC TCC AAA TGT GG-3

Fig 7B

SEQ. ID NO:10

ACA GTG ATG CAT TGG TTT GGG TTT TCA ACT TGG C

Fig 7C

SEQ. ID NO:11

ATA TAC ATG CAT CTG ACT TTC CCA TTG GGA CAT CTT TAA AG

Fig 7D

SEQ. ID NO:12

ACA GTG AGA TCT TTA GTG ATG GTG ATG GTG ATG AAT TCC ACA GCG AGG GGC
AGG GCT

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Fig. 8A
SEQ ID NO:13

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I	G	E	S	T	I	R	C	T	S	D	P	Q	G	N	G	V	W	S	S
P	A	P	R	C	G	I	H	H	H	H	H	H							

Fig. 8B

SEQ. ID NO: 14

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 CTCCCTGGGGAGCTGCCCGGCTGCTGCTGGTGCTGTTGTGCCTGCCGGCGTGTGGGGT
 GACTGTGGCCTTCCCCCAGATGTACCTAATGCCAGCCAGCTTGGAAAGGCCGTACAAGTTT
 CCCGAGGATACTGTAATAACGTACAAATGTGAAGAAAGCTTGTGAAAATTCTGGCGAGAAG
 GACTCAGTGATCTGCCTTAAGGGCAGTCAATGGTCAGATATTGAAGAGTTCTGCAATCGTAGC
 TGCGAGGTGCCAACAGGCTAAATTCTGCATCCCTCAAACAGCCTTATATCACTCAGAATTAT
 TTTCCAGTCGGTACTGTTGTGGAATATGAGTGCCGTCCAGGTTACAGAAGAGAACCTCTCTA
 TCACCAAAACTAACTTGCCCTCAGAATTAAAATGGTCCACAGCAGTCGAATTGTGAAATT
 AAATCATGCCCTAACCGGGAGAAATACGAAATGGTCAGATTGATGTACCAAGGTGGCATATTA
 TTTGGTGCAACCATCTCCTCTCATGTAACACAGGGTACAAATTATTGGCTCGACTTCTAGT
 TTTGTCTTATTCAGGCAGCTGTCCAGTGGAGTGACCCGTTGCCAGAGTGCAGAGAAATT
 TATTGTCCAGCACCAACAAATTGACAATGGAATAATTCAAGGGAACGTGACCATTATGGA
 TATAGACAGTCTGTAACGTATGCATGTAATAAAGGATTCAACATGATTGGAGAGCACTCTATT
 TATTGTACTGTGAATAATGATGAAGGAGAGTGGAGTGGCCCACCACTGAATGC
TCGAGTCCTAACAAATGCACGCCTCCAAATGTGGAAAATGGAATATTGGTATCTGACAAC
 AGAAGCTTATTTCTTAAATGAAGTTGTGGAGTTAGGTGTCAAGCCTGGCTTGTCA
 AAAGGACCCCGCCGTGTGAAGTGCCAGGCCCTGAACAAATGGGAGCCGGAGCTACCAAGC
 TGCTCCAGGGTATGTCAGCCACCTCCAGATGTCCTGCATGCTGAGCGTACCCAAAGGGAC
 AAGGACAACCTTCACCTGGCAGGAAGTGTCTACAGCTGTGAGGCCGCTACGACCTC
 AGAGGGCTCGCTATGCGCTGCACACCCAGGGAGACTGGAGGCCCTGCAGCCCCACA
 TGTGAAGTGAATCCTGTGATGACTTCATGGCCAACCTCTTAATGGCGTGTGCTATT
 CCAGTAAATCTCCAGCTGGAGCAAAAGTGGATTTGTTGTGATGAAGGATTCAATT
 AAAGGCAGCTCTGCTAGTTACTGTGTCTGGCTGGAATGGAAGCCTTGGAAATAGCAGT
 GTTCCAGTGTGTGAACAAATCTTGTCCAAGTCCTCCAGTATTCTTAATGGGAGACAC
 ACAGGAAAACCTCTGGAAGTCTTCCCTTGAAAAGCAGTAAATTACACATGCGACCCC
 CACCCAGACAGAGGGACGAGCTCGACCTCATTGGAGAGAGCACCATCCGCTGCACAAGT
 GACCCTCAAGGGAAATGGGTTGGAGCAGCCCTGCCCTCGCTGTGGAATTCTGGTCAC
 TGTCAAGCCCCAGATCATTCTGTTGCCAAGTTGAAAACCCAAACCAATGCATCTGAC
 TTTCCCATTGGACATCTTAAAGTACGAATGCCGTCTGAGTACTACGGGAGGCCATT
 TCTATCACATGTCTAGATAACCTGGTCTGGTCAAGTCCAAAGATGTCTGTAAACGTAAA
 TCATGTAAAACCTCCAGATCCAGTGAATGGCATGGTCATGTGATCACAGACATCCAG
GTTGGATCCAGAATCAACTATTCTGTACTACAGGGCACCGACTCATTGGTCACTCATCT
 GCTGAATGTATCCTCTCGGGCAATGCTGCCATTGGAGCACGAAGCCGCAATTGTCAA
 CGAATTCTGTGGCTACCCCCCACCATGCCAATGGAGATTCTTAATGACCAACAGA
 GAGAATTTCACTATGGATCAGTGGTGACCTACCGCTGCAATCCTGGAAGCGGAGGGAGA
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 GGCATCTGGAGCGGCCGGCCCTCAGTGCATTACACCTAACAAATGCACGCCTCAAAT
 GTGGAAAATGGAATTGGTATCTGACAACAGAAGCTTATTCTTCTAAATGAAGTTGTG
 GAGTTAGGTGTCAAGCCTGGCTTGTGATGAAAGGACCCCGCCGTGTGAAGTGCCAGGCC
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 GTCCTGCATGCTGAGCGTACCCAAAGGGACAAGGACAACCTTCAACCCGGCAGGAAGTG
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 CAGGGAGACTGGAGGCCCTGCAGCCCCACATGTGAAGTGAATCCTGTGATGACTTCATG
 GGCCAACTCTTAATGGCGTGTGCTATTCCAGTAAATCTCCAGCTGGAGCAAAAGTG

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 GCTGGAATGGAAAGCCTTGGAAATAGCAGTGTCCAGTGTGAACAAATCTTGTCCA
 AGTCCTCCAGTTATTCTTAATGGGAGACACACAGGAAAACCTCTGGAAGTCTTCCCTT
 GGAAAAGCAGTAAATTACACATGCGACCCCCACCCAGACAGAGGGACGAGCTCGACCTC
 ATTGGAGAGAGCACCATCCGCTGCACAAGTGACCCCTCAAGGGAATGGGGTTGGAGCAGC
 CCTGCCCTCGCTGGAAATTCATCACCATCACAAAGATCT

Fig. 9A
 SEQ ID NO:15

	M	T	V	A	R	P	S	V	P	A	A	L	P							
L	L	G	E	L	P	R	L	L	L	V	L	C	L	P	A	V	W	G		
D	C	G	L	P	P	D	V	P	N	A	Q	P	A	L	E	G	R	T	S	F
P	E	D	T	V	I	T	Y	K	C	E	E	S	F	V	K	I	P	G	N	K
D	S	V	I	C	L	K	G	S	Q	W	S	D	I	E	Y	F	C	T	Q	R
C	E	V	P	T	R	L	N	S	A	S	L	K	Q	P	Y	R	E	P	Q	S
F	P	V	G	T	V	V	E	Y	E	C	R	P	G	Y	R	F	P	C	G	L
S	P	K	L	T	C	L	Q	N	L	K	W	S	T	A	V	F	G	T	R	K
K	S	C	P	N	P	G	E	I	R	N	G	Q	I	D	V	F	G	S	C	I
F	G	A	T	I	S	F	S	S	C	N	T	G	Y	K	L	P	G	H	R	H
F	C	L	I	S	G	S	S	V	Q	W	S	D	P	L	F	P	E	R	E	I
Y	C	P	A	S	P	P	Q	I	D	N	G	I	F	Q	M	E	G	H	E	S
Y	R	Q	S	V	T	Y	A	C	N	K	G	I	F	T	P	I	P	G	C	I
Y	C	T	V	N	N	D	E	G	E	W	S	G	P	L	P	L	P	R	D	I
S	S	P	N	K	C	T	P	P	N	V	E	R	N	G	P	I	F	S	M	S
R	S	L	F	S	L	N	E	V	V	E	F	R	G	C	E	E	R	P	D	L
K	G	P	R	R	V	K	C	Q	A	L	N	K	W	A	E	E	Q	R	P	T
C	S	R	V	C	Q	P	P	P	D	V	F	Y	S	W	A	E	E	T	G	D
K	D	N	F	S	P	G	Q	Q	E	V	F	D	L	C	D	E	Q	R	P	L
R	G	A	A	S	M	R	C	T	P	Q	G	Q	L	W	S	G	R	G	Q	S
C	E	V	K	S	C	D	D	F	M	G	Q	L	W	C	D	E	L	P	G	N
P	V	N	L	Q	L	G	A	K	V	D	F	M	E	S	D	E	L	P	G	R
K	G	S	S	A	S	Y	C	V	L	A	G	M	P	V	I	N	Y	T	G	S
V	P	V	C	E	Q	I	F	C	P	F	G	P	A	V	N	T	Y	W	N	R
T	G	K	P	L	E	V	F	P	F	L	I	G	P	A	V	I	Y	Y	Y	T
H	P	D	R	G	T	S	F	D	L	I	G	P	R	S	T	C	I	G	T	R
D	P	Q	G	N	G	V	W	S	S	P	A	P	E	R	T	C	I	G	P	S
C	Q	A	P	D	H	F	L	F	A	K	R	K	P	T	Q	Y	T	Y	G	F
F	P	I	G	T	S	L	K	Y	E	C	R	L	P	E	K	D	Y	C	K	Q
S	I	T	C	L	D	N	L	V	W	S	S	P	P	K	D	V	I	T	G	R
S	C	K	T	P	P	D	P	V	N	G	M	V	H	V	H	R	L	I	H	S
V	G	S	R	I	N	Y	S	C	T	T	G	H	R	L	I	P	G	T	G	C
A	E	C	I	L	S	G	N	A	A	H	W	S	T	K	F	I	P	S	S	R
R	I	P	C	G	L	P	P	T	I	A	N	G	D	N	P	G	T	G	S	R
E	N	F	H	Y	G	S	V	V	T	Y	R	C	N	P	S	D	G	Q	P	R
K	V	F	E	L	V	G	E	P	S	I	Y	C	T	S	N	D	P	P	Q	V
G	I	W	S	G	P	A	P	Q	C	I	I	P	N	K	C	T	P	P	P	N

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V	E	N	G	I	L	V	S	D	N	R	S	L	F	S	L	N	E	V	V
E	F	R	C	Q	P	G	F	V	M	K	G	P	R	S	V	K	C	Q	A
L	N	K	W	A	E	P	E	P	S	C	S	R	V	C	Q	P	P	P	V
V	L	H	A	E	R	T	Q	R	D	K	D	N	F	S	S	M	C	D	V
F	Y	S	C	E	P	P	G	Y	D	L	R	G	A	S	S	C	G	R	P
Q	G	D	W	S	P	G	A	A	P	T	C	E	V	A	K	D	A	F	M
G	Q	L	L	N	G	R	V	L	F	P	V	N	L	Q	S	G	G	C	V
D	F	V	C	D	E	G	F	Q	L	K	G	S	S	C	E	A	F	F	M
A	G	M	E	S	L	W	N	S	S	V	P	V	C	P	E	Q	F	C	V
S	P	P	V	I	P	N	G	R	H	T	G	P	R	G	T	V	D	P	F
G	K	P	A	V	N	Y	T	C	D	P	H	P	D	Q	N	S	F	D	L
I	G	E	S	T	I	R	C	T	S	D	C	P	Q	A	G	H	F	S	A
P	A	P	R	T	C	G	I	L	G	H	D	F	P	I	T	S	L	K	E
K	L	K	T	Q	T	N	A	S	D	F	F	S	I	C	T	D	N	L	W
C	R	P	E	Y	Y	G	R	P	K	R	I	S	C	K	T	P	P	P	V
S	S	P	K	D	V	C	K	P	R	I	Q	V	H	R	L	I	G	H	S
G	M	V	H	V	I	T	D	D	I	T	T	G	H	S	T	K	P	S	Q
G	S	R	I	N	Y	S	C	T	A	A	H	W	R	T	K	G	H	I	R
A	E	C	I	L	S	G	N	A	T	I	A	N	S	D	F	S	T	S	N
R	I	P	C	G	L	P	P	P	T	T	Y	R	C	N	P	G	G	G	R
E	N	F	H	Y	G	S	V	V	T	S	I	Y	C	T	S	D	S	T	V
K	V	F	E	L	V	G	E	P	S	I	R	I	P	N	K	C	T	P	Q
G	I	W	S	G	P	A	P	Q	C	I	R	S	L	F	S	L	V	K	R
V	E	N	G	I	L	V	S	D	N	R	S	L	R	R	R	N	C	P	V
E	F	R	C	Q	P	G	F	V	M	K	G	P	R	R	R	Q	C	P	A
L	N	K	W	E	P	E	L	P	S	C	K	S	R	N	F	S	P	E	D
V	L	H	A	E	R	T	Q	R	D	K	R	G	A	K	S	C	P	E	V
F	Y	S	C	E	P	G	Y	D	L	T	C	E	V	N	L	Q	D	F	P
Q	G	D	W	S	P	G	A	A	P	T	F	V	S	S	C	G	R	M	V
G	Q	L	L	N	G	R	V	L	F	L	K	G	V	N	S	A	G	C	V
D	F	V	C	D	E	G	F	Q	L	P	K	V	P	V	C	E	Q	F	P
A	G	M	E	S	L	W	N	S	S	V	H	T	G	P	R	G	T	F	L
S	P	P	V	I	P	N	G	R	H	P	H	T	G	P	D	G	S	F	S
G	K	P	A	V	N	Y	T	C	D	P	H	P	H	Q	G	N	V	F	L
I	G	E	S	T	I	R	C	T	S	D	P	H	H	H	H	G	W	S	S
P	A	P	R	C	G	I	H	H	H	H	H	H	H	H	H	H	H	H	H

Fig 9B
SEQ. ID NO:16

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 CTCCTCGGGGAGCTGCCCGGCTGCTGCTGGTGCTGTTGTGCCTGCCGGCGTGTGGGGT
 GACTGTGGCCTTCCCCAGATGTACCTAATGCCAGCCAGCTTGTGAAAATTCCCTGGCGAGAAG
 CCCGAGGATACTGTAATAACGTACAAATGTGAAGAAAGCTTGTGAAAATTCCCTGGCGAGAAG
 GACTCAGTGATCTGCCTTAAGGGCAGTCATGGTCAGATATTGAAGAGTTCTGCAATCGTAGC
 TGCGAGGTGCCAACAGGCTAAATTCTGCATCCCTCAAACAGCCTATATCACTCAGAATTAT
 TTTCCAGTCGGTACTGTTGTGGAATATGAGTGCCGTCCAGGTTACAGAAGAGAACCTCTCTA

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TCACCAAAACTAACCTGCCTTCAGAATTAAAATGGTCCACAGCAGTCGAATTTGTAAAAAG
 AAATCATGCCCTAATCCGGGAGAAATACGAAATGGTCAGATTGATGTACCAGGTGGCATATTA
 TTTGGTGCAACCATCTCCTCTCATGTAACACAGGGTACAAATTATTGGCTCGACTCTAGT
 TTTTGTCTTATTCAGGCAGCTCTGTCCAGTGGAGTGACCCGTTGCCAGAGTGCAGAGAAATT
 TATTGTCCAGCACCACCAAAATTGACAATGGAATAATTCAAGGGAACGTGACCATTATGGA
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 ACAGGAAAACCTCTGGAAGTCTTCCCTTGGAAAAGCAGTAAATTACACATGCGACCCC
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 TGTCAAGCCCCAGATCATTTCTGTTGCCAGTTGAAAGTGCATCTGAC
 TTTCCCATTGGACATCTTAAAGTACGAATGCCGTCTGAGTACTACGGGAGGCCATT
 TCTATCACATGTCTAGATAACCTGGTCTGGTCAAGTCCAAAGATGTCTGTAAACGTAAA
 TCATGTAAAACCTCCAGATCCAGTGAATGGCATGGTCAAGTGTGATCACAGACATCCAG
 GTTGGATCCAGAATCAACTATTCTGTACTACAGGGCACCAGTCACTCATCT
 GCTGAATGTATCCTCTCGGGCAATGCTGCCATTGGAGCAGGAAGCCCAATTGTCAA
 CGAATTCTGTGGCTACCCCCCACCATCGCCAATGGAGATTCTTAGCACCAACAGA
 GAGAATTTCACATGGATCAGTGGTACCTACCGCTGCAATCCTGGAAGCGGGAGGGAGA
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 GGCAATGGTGCATGTGATCACAGACATCCAGGT
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 CGAATTCTGTGGCTACCCCCCACCATCGCCAATGGAGATTCTTAGCACCAACAGA
 GAGAATTTCACATGGATCAGTGGTGACCTACCGCTGCAATCCTGGAAGCGGGAGGGAGA

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AAGGTGTTGAGCTTGTGGGTGAGCCCTCCATATACTGCACCAAGCAATGACGATCAAGTG
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 CAGGGAGACTGGAGCCCTGCAGCCCCACATGTGAAGTGAAATCCTGTGATGACTTCATG
 GGCCAACTTCTTAATGGCCGTGTGCTATTCCAGTAAATCTCCAGCTGGAGCAAAAGTG
 GATTGTTGTTGTGATGAAGGATTCAATTAAAAGGCAGCTCTGCTAGTTATTGTGTCTTG
 GCTGGAATGGAAAGCCTTGGAAATAGCAGTGTGCTGAAACAAATCTTTGTCCA
 AGTCCTCCAGTTATTCTTAATGGGAGACACACAGGAAACCTCTGGAAGTCTTCCCTT
 GGAAAAGCAGTAAATTACACATGCGACCCCCACCCAGACAGAGGGACGAGCTCGACCTC
 ATTGGAGAGAGCACCATCCGCTGCACAAGTGACCCCTCAAGGGAATGGGTTGGAGCAGC
 CCTGCCCTCGCTGTCCAATTCACCATCACCATCACTAAAGATCT

Fig. 10A

SEQ. ID NO:17

ATA TAC GAA TTC TGG TTG AGT CCA AAT ATG GTC CC

Fig. 10B

SEQ. ID NO:18

ACA GTG AGA TCT TTA TCA TTT ACC CGG AGA CAG GGA G

Fig. 11A

SEQ. ID NO:19

	M	T	V	A	R	P	S	V	P	A	A	L	P							
L	L	G	E	L	P	R	L	L	V	L	L	C	L	P	A	V	W	G		
D	C	G	L	P	P	D	V	P	N	A	Q	P	A	L	E	G	R	T	S	F
P	E	D	T	V	I	T	Y	K	C	E	E	S	F	V	K	I	P	G	E	K
D	S	V	I	C	L	K	G	S	Q	W	S	D	I	E	E	F	C	N	R	S
C	E	V	P	T	R	L	N	S	A	S	L	K	Q	P	Y	I	T	Q	N	Y
F	P	V	G	T	V	V	E	Y	E	C	R	P	G	Y	R	R	E	P	S	L
S	P	K	L	T	C	L	Q	N	L	K	W	S	T	A	V	E	F	C	K	K
K	S	C	P	N	P	G	E	I	R	N	G	Q	I	D	V	P	G	G	I	L
F	G	A	T	I	S	F	S	C	N	T	G	Y	K	L	F	G	S	T	S	S
F	C	L	I	S	G	S	S	V	Q	W	S	D	P	L	P	E	C	R	E	I
Y	C	P	A	P	P	Q	I	D	N	G	I	I	Q	G	E	R	D	H	Y	G
Y	R	Q	S	V	T	Y	A	C	N	K	G	F	T	M	I	G	E	H	S	I
Y	C	T	V	N	N	D	E	G	E	W	S	G	P	P	P	E	<u>C</u>			
S	S	P	N	K	C	T	P	P	N	V	E	N	G	I	L	V	<u>S</u>	D	N	

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R	S	L	F	S	L	N	E	V	E	F	R	C	Q	P	G	F	V	M	
K	G	P	R	R	V	K	C	Q	A	L	N	K	W	E	P	E	L	P	S
C	S	R	V	C	Q	P	P	D	V	L	H	A	E	R	T	Q	R	D	
K	D	N	F	S	P	G	Q	E	V	F	Y	S	C	E	P	G	Y	D	L
R	G	A	A	S	M	R	C	T	P	Q	G	D	W	S	P	A	A	P	T
C	E	V	K	S	C	D	D	F	M	G	Q	L	L	N	G	R	V	L	F
P	V	N	L	Q	L	G	A	K	V	D	F	V	C	D	E	G	F	Q	L
K	G	S	S	A	S	Y	C	V	L	A	G	M	E	S	L	W	N	S	S
V	P	V	C	E	Q	I	F	C	P	S	P	P	V	I	P	N	G	R	H
T	G	K	P	L	E	V	F	P	F	G	K	A	V	N	Y	T	C	D	P
H	P	D	R	G	T	S	F	D	L	I	G	E	S	T	I	R	C	T	S
D	P	Q	G	N	G	V	W	S	S	P	A	P	R	C	G	I	L		
V	E	S	K	Y	G	P	P	C	P	S	C	P	A	P	E	F	L		
G	G	P	S	V	F	L	F	P	P	K	P	K	D	T	L	M	I	S	R
T	P	E	V	T	C	V	V	V	D	V	S	Q	E	D	P	E	V	Q	F
N	W	Y	V	D	G	V	E	V	H	N	A	K	T	K	P	R	E	E	Q
F	N	S	T	Y	R	V	V	S	V	L	T	V	L	H	Q	D	W	L	N
G	K	E	Y	K	C	K	V	S	N	K	G	L	P	S	S	I	E	K	T
I	S	K	A	K	G	Q	P	R	E	P	Q	V	Y	T	L	P	P	S	Q
E	E	M	T	K	N	Q	V	S	L	T	C	L	V	K	G	F	Y	P	S
D	I	A	V	E	W	E	S	N	G	Q	P	E	D	N	Y	K	T	T	P
P	V	L	D	S	D	G	S	F	F	L	Y	S	R	L	T	V	D	K	S
R	W	Q	E	G	N	V	F	S	C	S	V	M	H	E	A	L	H	N	H
Y	T	Q	K	S	L	S	L	S	P	G	K								

Fig. 11B
SEQ. ID NO:20

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 CTCCTCGGGGAGCTGCCCCGGCTGCTGCTGGTGCTGTTGTGCCTGCCGGCGTGTGGGGT
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 TTTCCAGTCGGTACTGTTGTGGAATATGAGTGCCGTCCAGGTTACAGAAGAGAACCTCTCTA
 TCACCAAAACTAACTTGCCTTCAGAATTAAAATGGTCCACAGCAGTCGAATTGTAAAAG
 AAATCATGCCCTAATCCGGGAGAAATCGAAATGGTCAGATTGATGTACCAAGGTGGCATATTA
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 TTTTGTCTTATTCAGGCAGCTGTCCAGTGGAGTGACCCGTTGCCAGAGTGCAGAGAAATT
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 TATTGTACTGTGAATAATGATGAAGGAGAGTGGAGTGGCCCACCACTGAATGC
TCGAGTCCTAACAAATGCACGCCCTCAAATGTGGAAAATGGAATTGTTATCTGACAAAC
 AGAAGCTTATTTCTTAAATGAAGTTGTGGAGTTAGGTGTCAAGCCTGGCTTGTCA
 AAAGGACCCCGCCGTGTGAAGTGCCAGGCCCTGAACAAATGGGAGGCCGGAGCTACCAAGC
 TGCTCCAGGGTATGTCAGGCCACCTCCAGATGTCCTGCATGCTGAGCGTACCCAAAGGGAC
 AAGGACAACTTTCACCTGGGCAGGAAGTGTCTACAGCTGTGAGCCGGCTACGACCTC

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AGAGGGGCTGCGTCTATGCGCTGCACACCCCCAGGGAGACTGGAGGCCCTGCAGCCCCACA
TGTGAAGTGAAATCCTGTGATGACTTCATGGGCCAACTTCTTAATGGCCGTGTGCTATT
CCAGTAAATCTCCAGCTTGGAGCAAAAGTGGATTGTGATGAAGGATTCAATTAA
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ACAGGAAAACCTCTGGAAGTCTTCCCTTGGAAAAGCAGTAAATTACACATGCGACCCC
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GACCCTCAAGGGAATGGGTTGGAGCAGCCCTGCCCTCGCTGTGGATTCTG
GTTGAGTCAAATATGGTCCCCCATGCCCATCATGCCAGCACCTGAGTTCTG
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AACTGGTACGTGGATGGCGTGGAGGTGCATAATGCCAAGACAAAGCCGGAGGAGCAG
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CCCGTGCCTGGACTCCGACGGCTTCTACAGCAGGCTAACCGTGGACAAGAGC
AGGTGGCAGGAGGGAAATGTCTTCTCATGCTCCGTGATGCATGAGGCTCTGCACAAACCAC
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Fig. 12A

SEQ. ID NO:21

ATA TAC GAA TTC TGG GTC ACT GTG AGG AGC CAC CAA CAT TTG AAG C

Fig. 12B

SEQ. ID NO:22

ACÀ GTG AGA TCT TTA GTG ATG GTG ATG GTG ATG CGA CAC TTT AAG ACA CTT
TGG AAC

Fig. 13A

SEQ. ID NO:23

M	T	V	A	R	P	S	V	P	A	A	L	P	L	C	L	P	A	V	W	G	F	K	S	Y	L	H			
L	L	G	E	L	P	R	L	P	N	C	E	Q	S	A	F	V	E	K	R	T	G	N	O	P	C	G	T	R	
D	C	D	T	V	I	C	L	K	G	S	Y	E	S	Q	H	E	P	Y	R	E	F	C	T	E	F	G	S	C	
P	S	E	V	P	T	R	V	L	N	S	Y	E	W	S	K	Q	G	T	V	E	P	G	H	F	H	R	E	P	G
K	F	S	V	G	T	V	V	E	Q	N	I	R	C	K	W	G	T	H	D	V	F	G	C	T	E	F	G	T	R
S	K	C	K	L	T	C	C	Q	E	N	I	R	S	N	W	G	Y	K	L	F	P	G	H	F	H	R	E	P	G
G	F	C	C	A	T	H	S	G	F	S	S	C	N	T	W	S	D	P	L	L	P	G	H	F	H	R	E	P	G
F	F	C	C	L	H	S	G	S	S	V	Q	W	W	T	S	S	P	L	E	E	P	G	H	F	H	R	E	P	G

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Y	C	P	A	P	P	Q	I	D	N	G	I	I	Q	G	E	R	D	H	Y	G
Y	R	Q	S	V	T	Y	A	C	N	K	G	F	T	M	I	G	E	H	S	I
Y	C	T	V	N	N	D	E	G	E	W	S	G	P	P	P	E	C	S	D	N
S	S	P	N	K	C	T	P	P	N	V	E	N	G	I	L	V	G	F	V	M
R	S	L	F	S	L	N	E	V	V	E	F	R	C	Q	P	G	F	D	V	N
K	G	P	R	R	V	K	C	Q	A	L	N	K	W	E	P	E	L	P	S	
C	S	R	V	C	Q	P	P	P	D	V	L	H	A	E	R	T	Q	R	D	
K	D	N	F	S	P	G	Q	E	V	F	Y	S	C	E	P	G	Y	D	L	
R	G	A	A	S	M	R	C	T	P	Q	G	D	W	S	P	A	A	P	T	
C	E	V	K	S	C	D	D	F	M	G	Q	L	L	N	G	R	V	L	F	
P	V	N	L	Q	L	G	A	K	V	D	F	V	C	D	E	G	F	Q	L	
K	G	S	S	A	S	Y	C	V	L	A	G	M	E	S	L	W	N	S	S	
V	P	V	C	E	Q	I	F	C	P	S	P	P	V	I	P	N	G	R	H	
T	G	K	P	L	E	V	F	P	F	G	K	A	V	N	Y	T	C	D	P	
H	P	D	R	G	T	S	F	D	L	I	G	E	S	T	I	R	C	T	S	
D	P	Q	G	N	G	V	W	S	S	P	A	P	R	C	G	I	L	G	H	
C	E	E	P	P	T	F	E	A	M	E	L	I	G	K	P	K	P	Y	Y	
E	I	G	E	R	V	D	Y	K	C	K	K	G	Y	F	Y	I	P	P	L	
A	T	H	T	I	C	D	R	N	H	T	W	L	P	V	S	D	D	A	C	
Y	R	E	T	C	P	Y	I	R	D	P	L	N	G	Q	A	V	P	A	N	
G	T	Y	E	F	G	Y	Q	M	H	F	I	C	N	E	G	Y	Y	L	I	
G	E	E	I	L	Y	C	E	L	K	G	S	V	A	I	W	S	G	K	P	
P	I	C	E	K	V	L	C	T	P	P	K	I	K	N	G	K	H	T	A	
F	S	E	V	E	V	F	E	Y	L	D	A	V	T	Y	S	C	D	P	N	
P	G	P	D	P	F	S	L	I	G	E	S	T	I	Y	C	G	D	N	S	
V	W	S	R	R	A	A	P	E	C	K	V	V	K	C	R	F	P	V	V	
N	G	K	Q	I	S	G	F	G	K	K	F	Y	Y	K	A	T	V	M	F	
E	C	D	K	G	F	Y	L	D	G	S	D	T	I	V	C	D	S	N	S	
T	W	D	P	P	V	P	K	C	L	K		V	S	H	H	H	H	H	H	

Fig. 13B

SEQ. ID NO:24

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 CTCCTCGGGGAGCTGCCCCGGCTGCTGCTGCTGGCTGCTGCTGTTGTGCCTGCCGGCGTGTGGGGT
 GACTGTGGCCTTCCCCAGATGTACCTAATGCCAGCCAGCTTGGAAAGGCCGTACAAGTTTT
 CCCGAGGATACTGTAATAACGTACAAATGTGAAGAAAGCTTGTGAAAATTCCCTGGCGAGAAAG
 GACTCAGTGTACTGCCTTAAGGGCAGTCAATGGTCAGATATTGAAGAGTTCTGCAATCGTAGC
 TGCGAGGTGCCAACAGGCTAAATTCTGCATCCCTCAAACAGCCTTATATCACTCAGAATTAT
 TTTCCAGTCGGTACTGTTGGAAATATGAGTGCCGTCCAGGTTACAGAAGAGAACCTCTCTA
 TCACCAAAACTAACTTGCCCTCAGAATTAAATGGTCCACAGCAGTCGAATTGTGAACTTGTAAAAAG
 AAATCATGCCCTAATCCGGGAGAAATCGAAATGGTCAGATTGATGTACCAGGTGGCATATTA
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 TATTGTCCAGCACCACCAAAATTGACAATGGAATAATTCAAGGGAAACGTGACCATTATGGA
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 AGAAGCTTATTTCCTTAAATGAAGTTGTGGAGTTAGGTGTAGCCTGGCTTGTATG

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AAAGGACCCCGCCGTGTGAAGTGCCAGGCCCTGAACAAATGGGAGCCGGAGCTACCAAGC
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ACAGGAAAACCTCTGGAAAGTCTTCCCTTGGAAAAGCAGTAAATTACACATGCGACCCC
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GGTGAAGAAATTCTATATTGTGAACCTAAAGGATCAGTAGCAATTGGAGCGGTAAGCCC
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GTGTGGAGTCGTGCTCCAGAGTGTAAAGTGGTCAAATGTCGATTCCAGTAGTCGAA
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GAATGCGATAAGGGTTTACCTCGATGGCAGCGACACAATTGTCTGTGACAGTAACAGT
ACTTGGGATCCCCCAGTTCCAAAGTGTCTAAA//GTGTCG//CATCACCATCAC
TAAAGATCT

WESTERN BLOT OF HYBRID PROTEINS
DAF-IgG4, DAF-CR1BB, and DAF-CR1B

Anti-DAF
IA10 mAb

Anti-CR1
E11 mAb

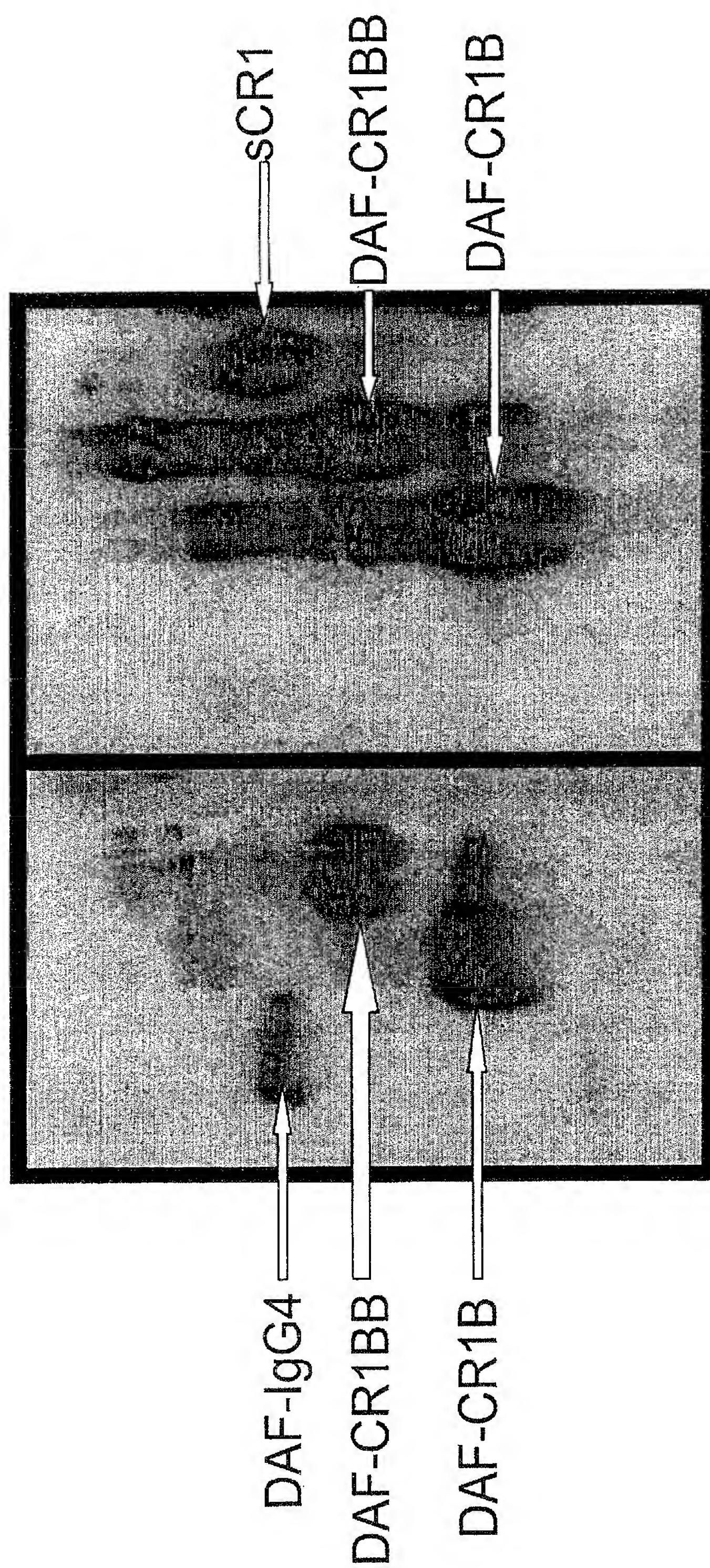


Fig. 14

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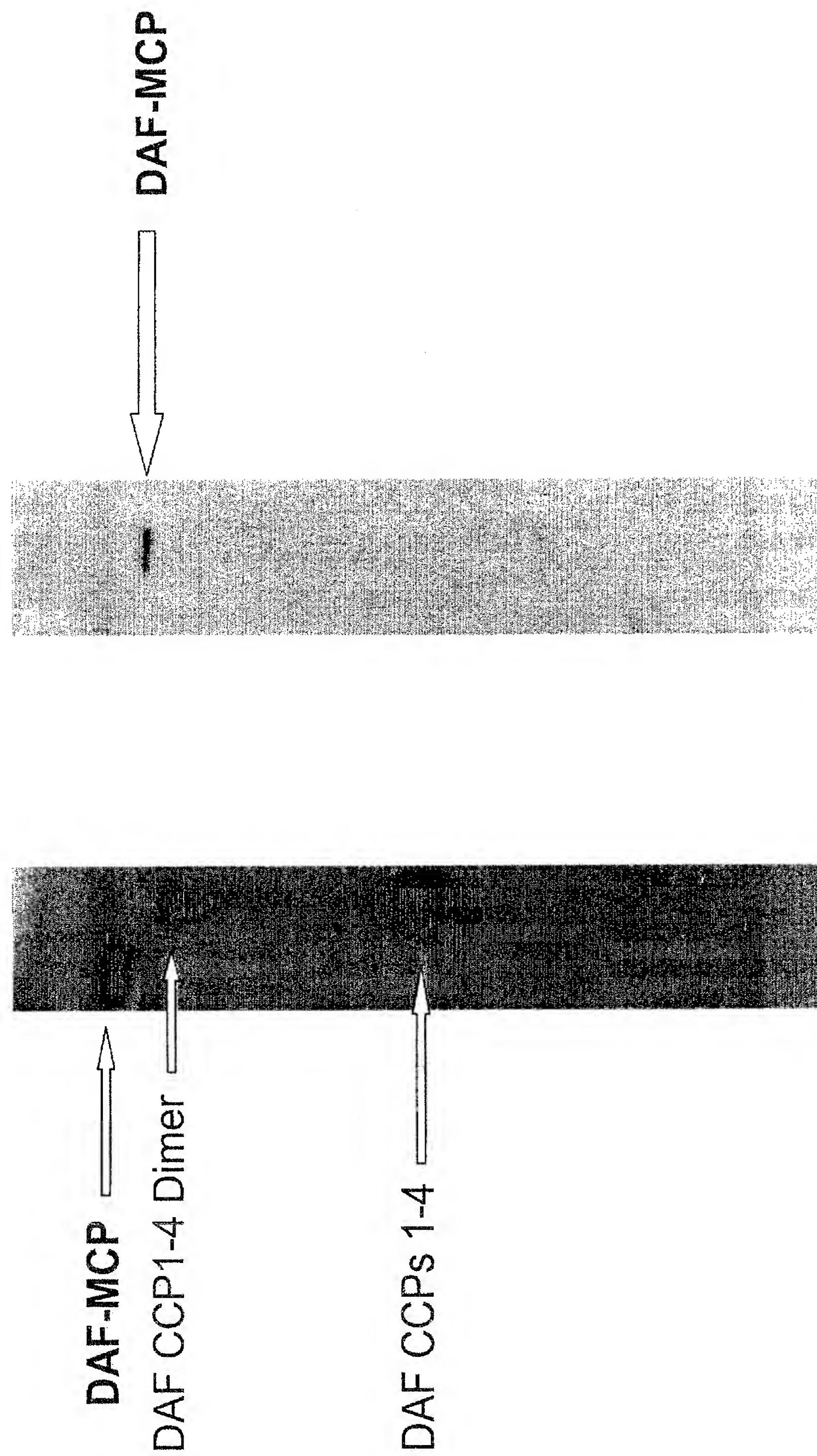
Western Blot of DAF-MCP

Anti-DAF

IA10

Anti-MCP

GB24

**Fig. 15**

Whole Serum Hemolytic Assay
DAF-CR1BB vs. Soluble CR1

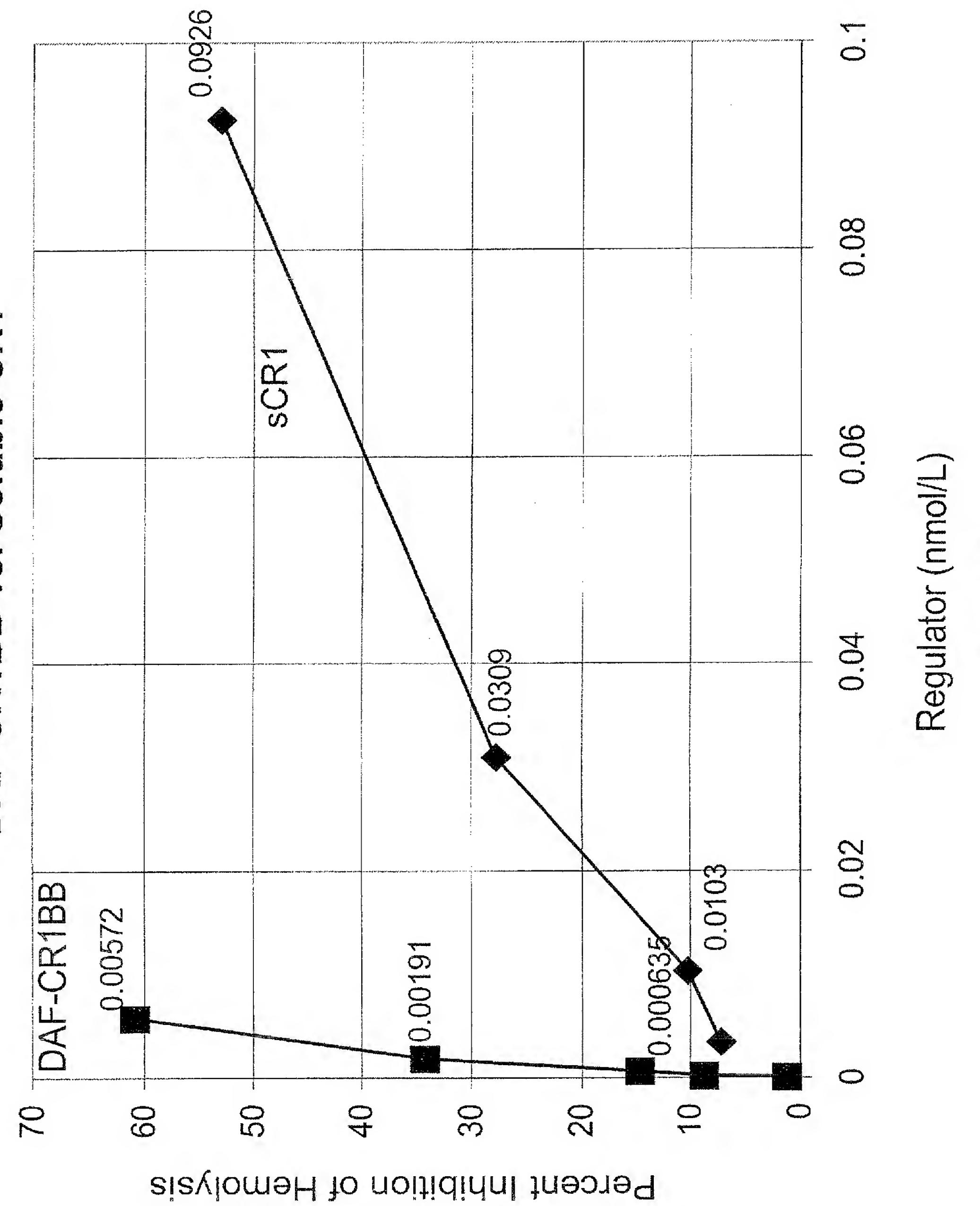


Fig. 16

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Whole Serum Hemolytic Assay
DAF-MCP Hybrid vs. DAF CCPs 1-4

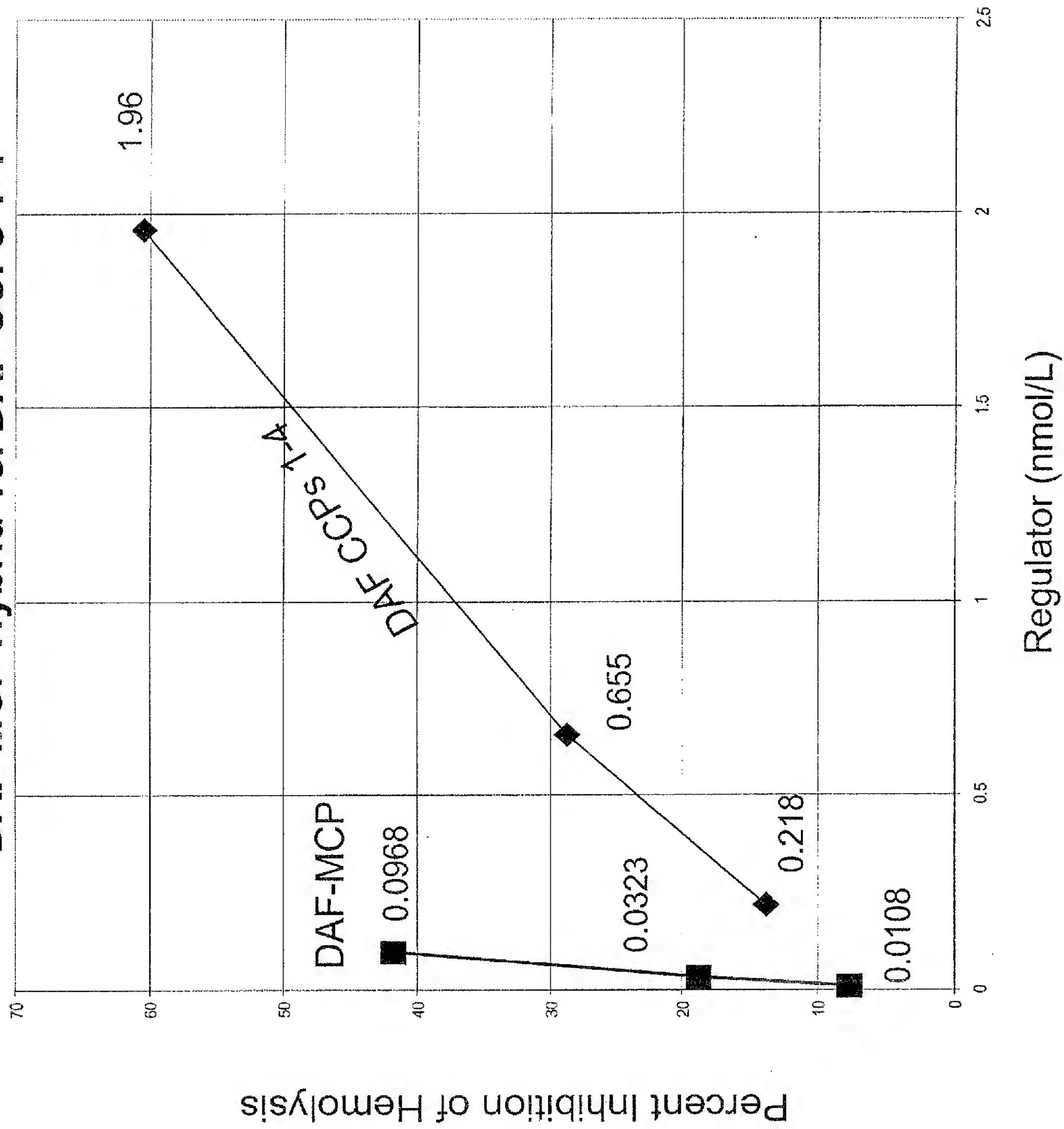


Fig. 17

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Classical Pathway C3 Convertase Decay

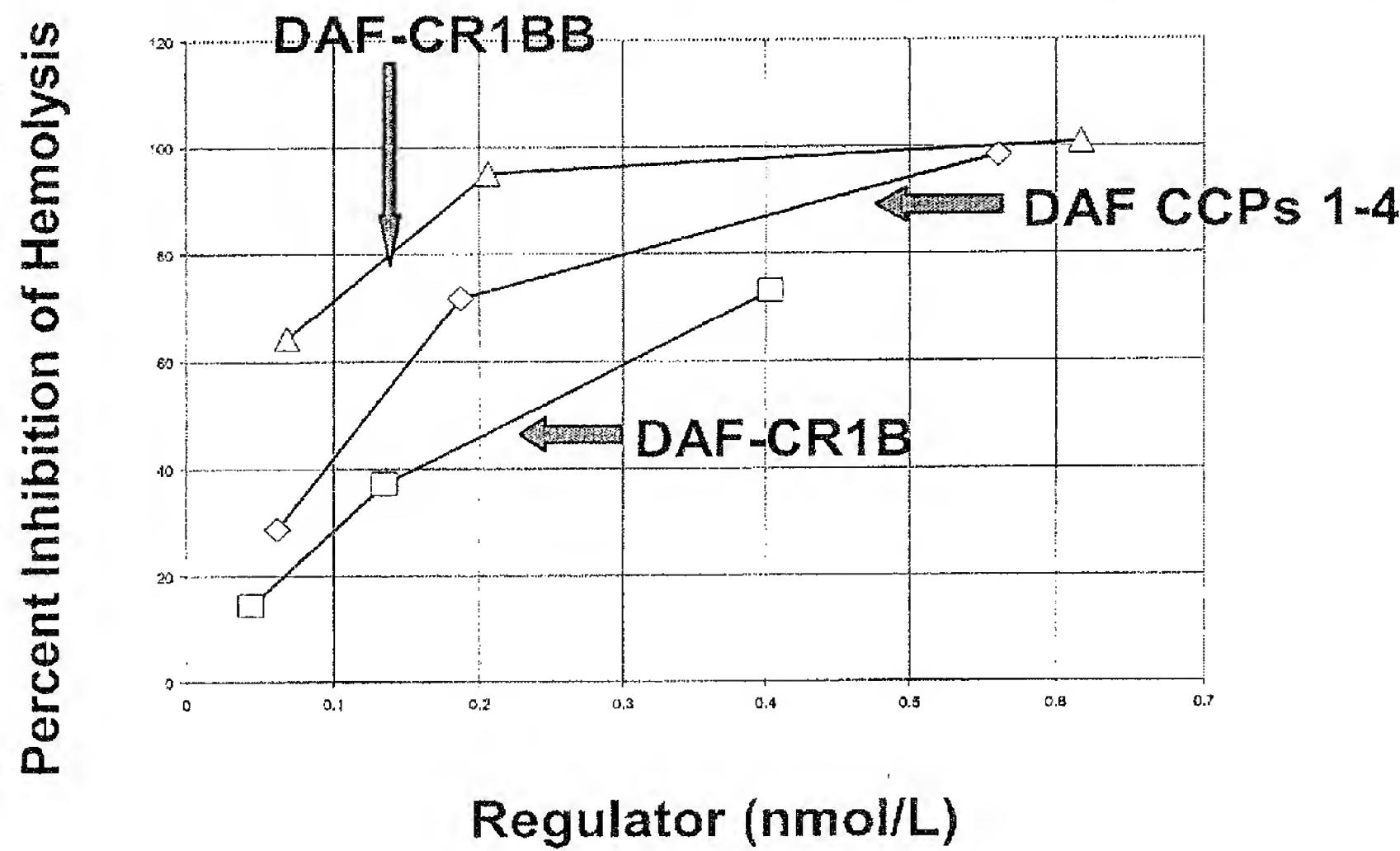


Fig. 18A

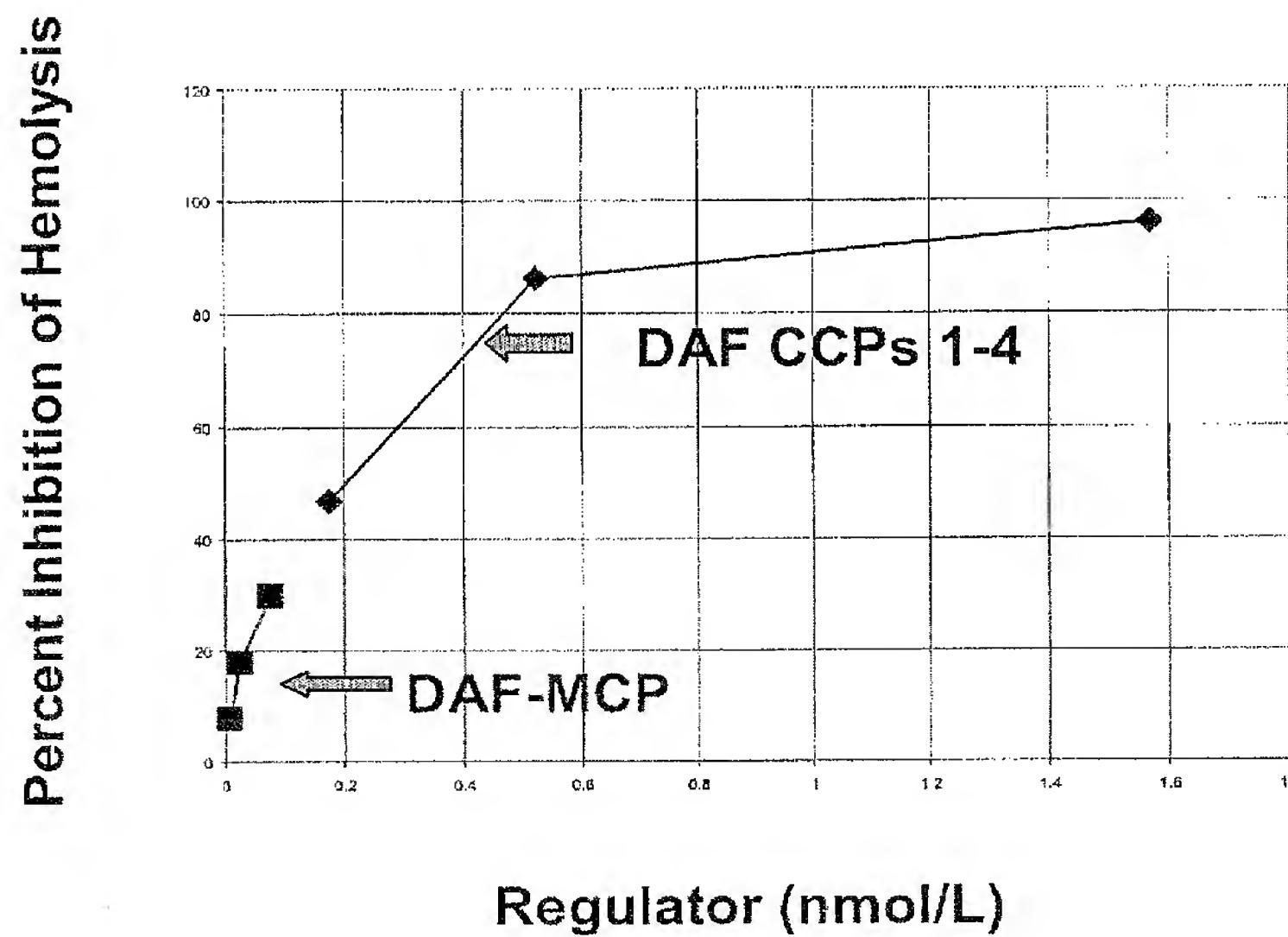
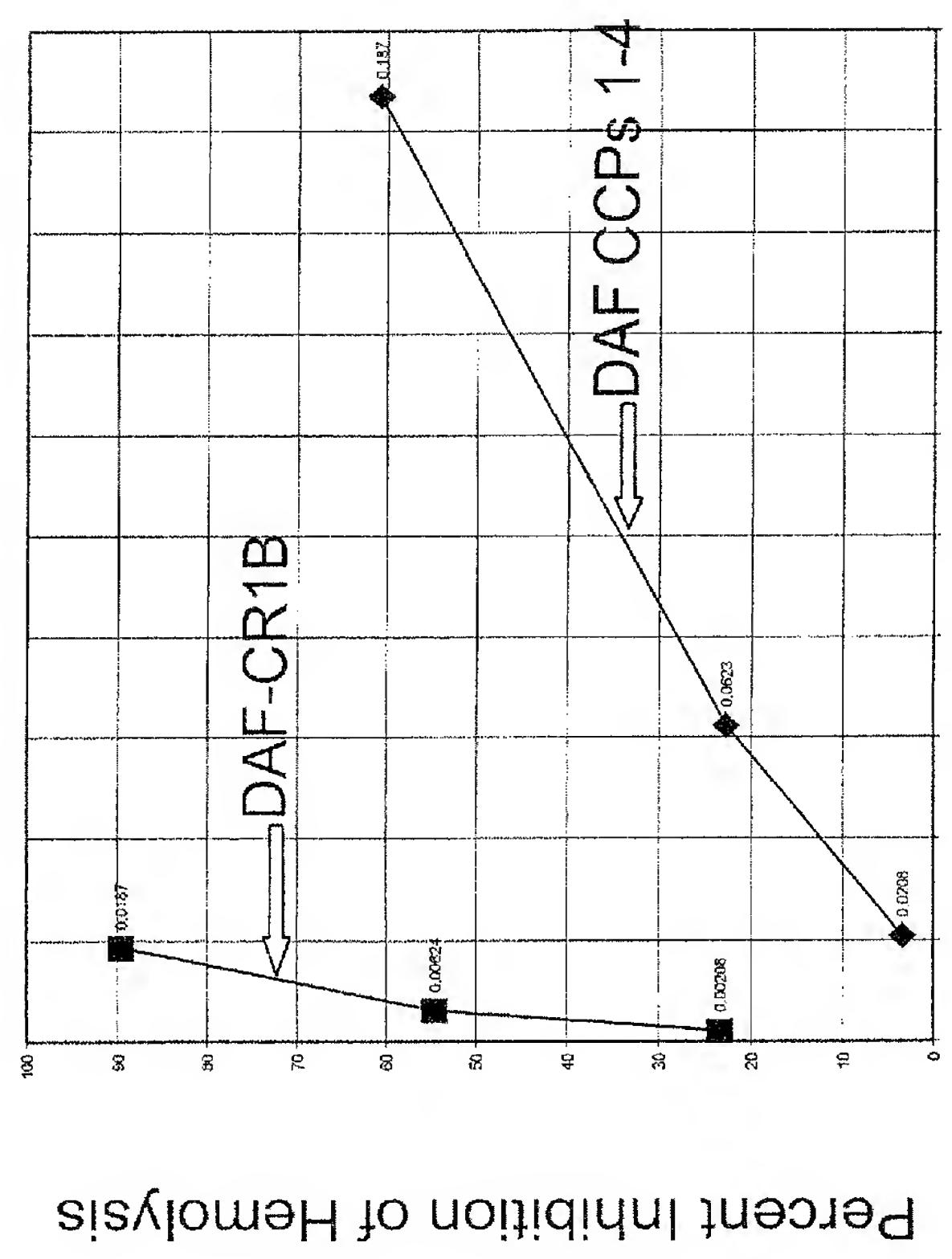


Fig. 18B

Classical Pathway C5 Convertase Decay
DAF-CR1B vs DAF CCPs 1-4



Regulator (nmol/L)

Fig. 19

Classical Pathway C5 Convertase Decay DAF-CR1BB vs sCR1 vs DAF-CR1B

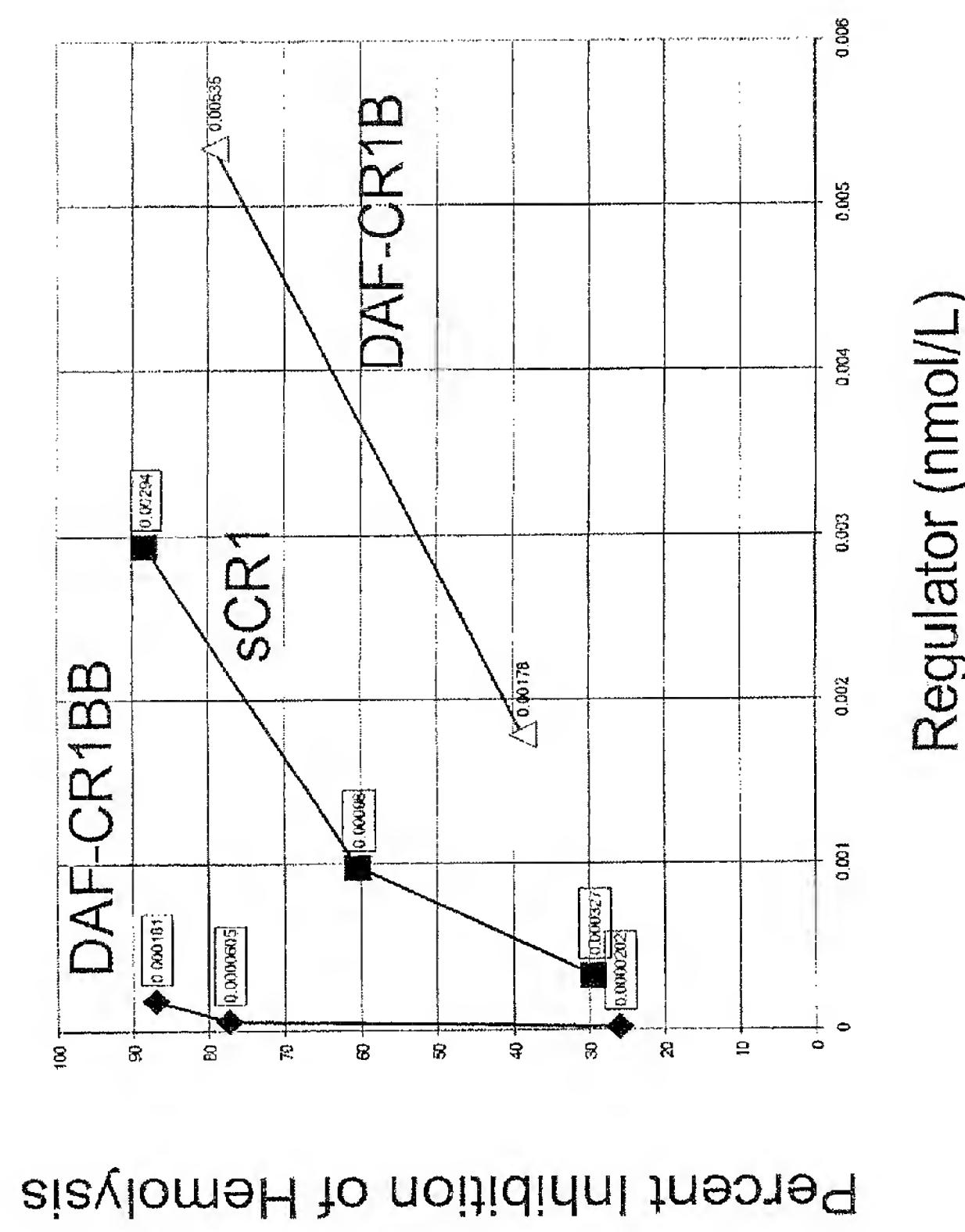


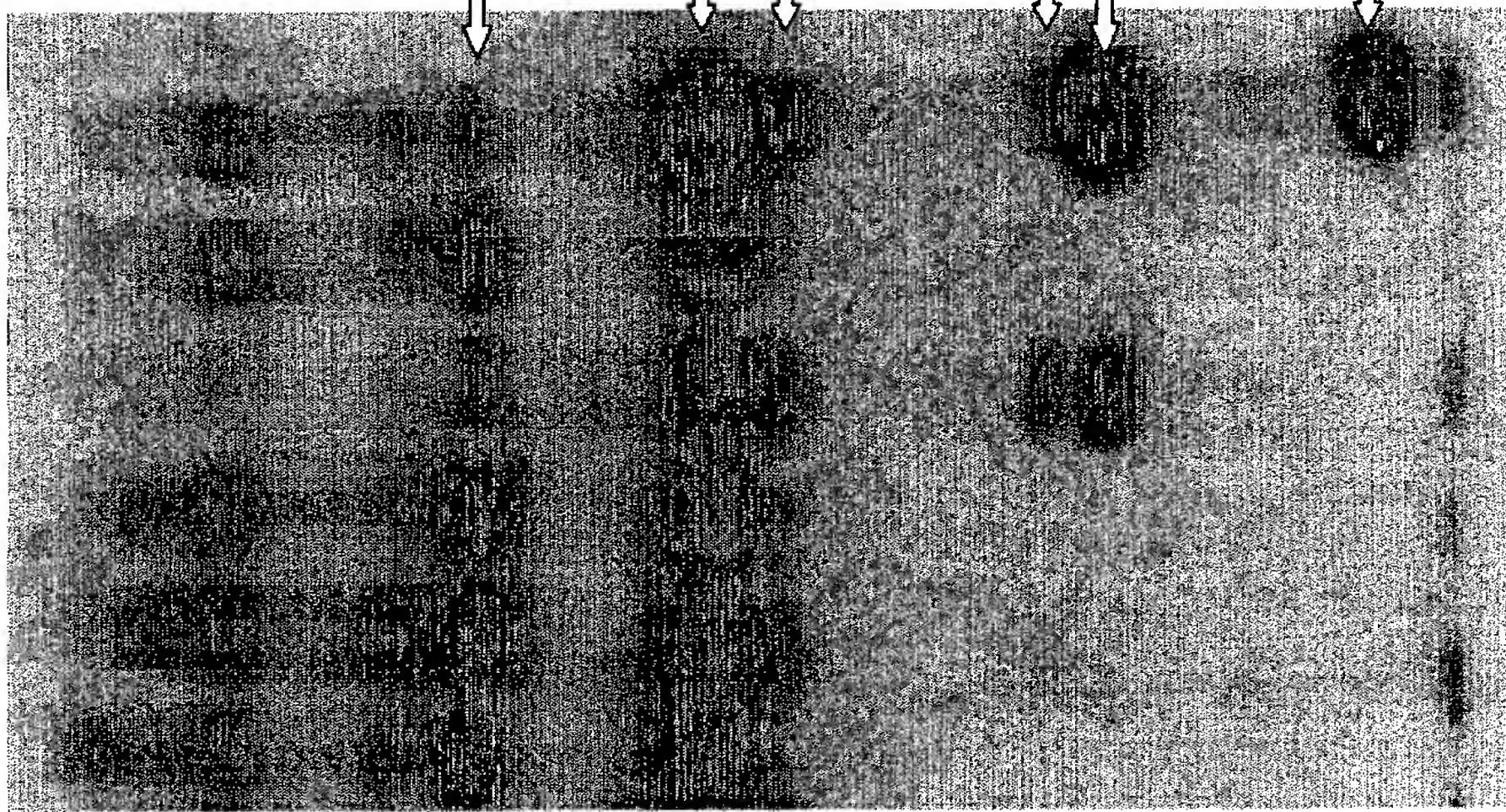
Fig. 20

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Cell-bound ($E_{sh}C4b3b$) Cofactor Assay

Cell Supernatant
(Anti-human C3 pAb)
COS SN DAF-MCP DAF-CR1BB

+| +| +|

**Fig. 21**

Cell-bound (EshC4b3b) Cofactor Assays Cell Supernatant

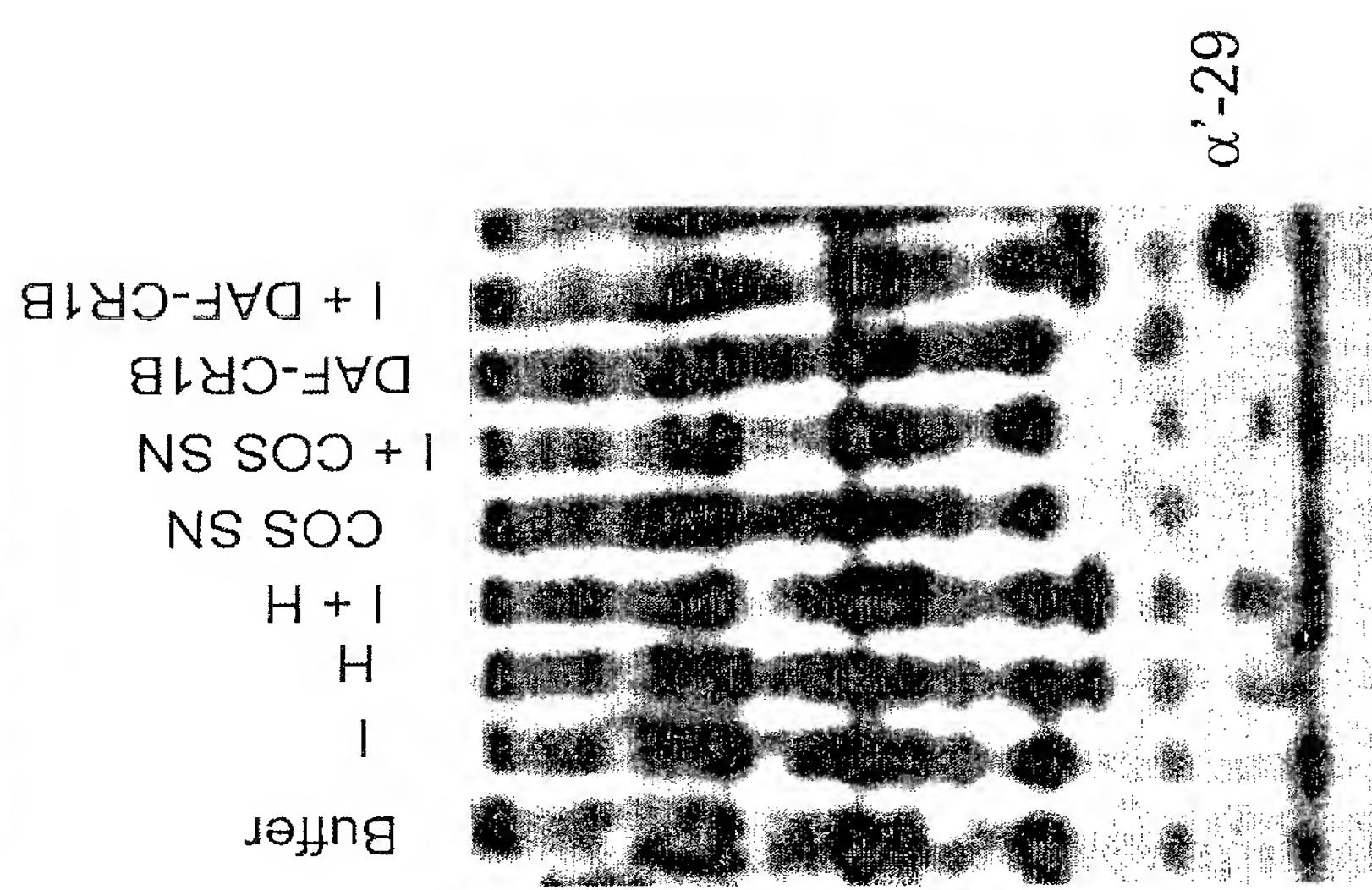


Fig. 22